## **RAW SEQUENCE LISTING**

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) no errors detected.

Application Serial Number:	_09	/820	,33	9B	
Source:			IFU	116	
Date Processed by STIC:		12	//3	105	

## ENTERED



## IFW16

RAW SEQUENCE LISTING DATE: 12/13/2005
PATENT APPLICATION: US/09/820,339B TIME: 14:01:37

Input Set : A:\sequence listing.txt

Output Set: N:\CRF4\12132005\1820339B.raw

```
3 <110> APPLICANT: FUCHS, Sara
              BARCHAN, Dora
      5
              SOUROUJON, Miriam C.
      7 <120> TITLE OF INVENTION: RECOMBINANT FRAGMENTS OF THE HUMAN ACETYLCHOLINE RECEPTOR
AND
              THEIR USE FOR TREATMENT OF MYASTHENIA GRAVIS
     10 <130> FILE REFERENCE: FUCHS=2A
C--> 12 <140> CURRENT APPLICATION NUMBER: US/09/820,339B
C--> 12 <141> CURRENT FILING DATE: 2001-03-29
     12 <150> PRIOR APPLICATION NUMBER: 09/423,398
     13 <151> PRIOR FILING DATE: 1999-11-08
     15 <150> PRIOR APPLICATION NUMBER: PCT/IL98/00211
     16 <151> PRIOR FILING DATE: 1998-05-06
     18 <160> NUMBER OF SEO ID NOS: 32
     20 <170> SOFTWARE: PatentIn version 3.3
     22 <210> SEQ ID NO: 1
     23 <211> LENGTH: 630
     24 <212> TYPE: DNA
     25 <213> ORGANISM: Homo sapiens
     27 <400> SEQUENCE: 1
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     30 ccagtggaag accaccgcca ggtcgtggag gtcaccgtgg gcctgcagct gatacagctc
                                                                              120
     32 atcaatgtgg atgaagtaaa tcagatcgtg acaaccaatg tgcgtctgaa acagcaatgg
                                                                              180
     34 gtggattaca acctaaaatg gaatccagat gactatggcg gtgtgaaaaa aattcacatt
                                                                              240
     36 ccttcagaaa agatctggcg cccagacctt gttctctata acgatgcaga tggtgacttt
                                                                               300
     38 gctattgtca agttcaccaa agtgctcctg cagtacactg gccacatcac gtggacacct
                                                                              360
     40 ccagccatct ttaaaagcta ctgtgagatc atcgtcaccc actttccctt tgatgaacag
                                                                              420
     42 aactgcagca tgaagctggg cacctggacc tacgacggct ctgtcgtggc catcaacccg
                                                                              480
     44 qaaaqcqacc aqccaqacct qaqcaacttc atqqaqaqcq qqqaqtqqqt qatcaaqqaq
                                                                              540
     46 tecegggget ggaageacte egtgaeetat teetgetgee eegacaeeee etacetggae
                                                                              600
                                                                              630
     48 atcacctacc acttcgtcat gcagcgcctg
     51 <210> SEQ ID NO: 2
     52 <211> LENGTH: 210
     53 <212> TYPE: PRT
     54 <213> ORGANISM: Homo sapiens
     56 <400> SEQUENCE: 2
     58 Ser Glu His Glu Thr Arg Leu Val Ala Lys Leu Phe Lys Asp Tyr Ser
     59 1
                                             10
                                                                 15
     62 Ser Val Val Arg Pro Val Glu Asp His Arg Gln Val Val Glu Val Thr
     63
                    20
                                         25
                                                             30
     66 Val Gly Leu Gln Leu Ile Gln Leu Ile Asn Val Asp Glu Val Asn Gln
                                    40
     70 Ile Val Thr Thr Asn Val Arg Leu Lys Gln Gln Trp Val Asp Tyr Asn
     71
                                55
            50
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Input Set : A:\sequence listing.txt
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74 Leu Lys Trp Asn Pro Asp Asp Tyr Gly Gly Val Lys Lys Ile His Ile				
75 65 70 75 80				
78 Pro Ser Glu Lys Ile Trp Arg Pro Asp Leu Val Leu Tyr Asn Asn Ala				
79 85 90 95				
82 Asp Gly Asp Phe Ala Ile Val Lys Phe Thr Lys Val Leu Leu Gln Tyr				
83 100 105 110				
86 Thr Gly His Ile Thr Trp Thr Pro Pro Ala Ile Phe Lys Ser Tyr Cys				
87 115 120 125				
90 Glu Ile Ile Val Thr His Phe Pro Phe Asp Glu Gln Asn Cys Ser Met				
91 130 135 140				
94 Lys Leu Gly Thr Trp Thr Tyr Asp Gly Ser Val Val Ala Ile Asn Pro				
95 145 · 150 155 160				
98 Glu Ser Asp Gln Pro Asp Leu Ser Asn Phe Met Glu Ser Gly Glu Trp				
99 165 170 175				
102 Val Ile Lys Glu Ser Arg Gly Trp Lys His Ser Val Thr Tyr Ser Cys				
103 180 185 190				
106 Cys Pro Asp Thr Pro Tyr Leu Asp Ile Thr Tyr His Phe Val Met Gln				
107 195 200 205				
110 Arg Leu				
111 210				
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116 <212> TYPE: DNA				
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122 catctgcagg atgag	75			
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126 <211> LENGTH: 25				
127 <212> TYPE: PRT				
128 <213> ORGANISM: Homo sapiens				
130 <400> SEQUENCE: 4				
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133 1 5 10 15				
136 Pro Leu Phe Ser His Leu Gln Asp Glu				
137 20 25				
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141 <211> LENGTH: 705				
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150 atcaatgtgg atgaagtaaa tcagatcgtg acaaccaatg tgcgtctgaa acagggtgac	180			
150 atcaatgtgg atgaagtaaa tcagatcgtg acaaccaatg tgcgtctgaa acagggtgac 152 atggtagatc tgccacgccc cagctgcgtg actttgggag ttcctttgtt ttctcatctg				
152 atggtagatc tgccacgccc cagctgcgtg actttgggag ttcctttgtt ttctcatctg	180			
	180 240			
152 atggtagatc tgccacgccc cagctgcgtg actttgggag ttcctttgtt ttctcatctg 154 caggatgagc aatgggtgga ttacaaccta aaatggaatc cagatgacta tggcggtgtg	180 240 300			
152 atggtagate tgecaegeee cagetgegtg aetttgggag tteetttgtt tteteatetg 154 caggatgage aatgggtgga ttacaaeeta aaatggaate cagatgaeta tggeggtgtg 156 aaaaaaatte acatteette agaaaagate tggegeeeag aeettgttet etataaegat	180 240 300 360			

Input Set : A:\sequence listing.txt
Output Set: N:\CRF4\12132005\I820339B.raw

	ccctttgatg aacagaactg cagcatgaag ctgggcacct ggacctacga cggctctgt			
	gtggccatca acccggaaag cgaccagcca gacctgagca acttcatgga gagcgggga			
	tgggtgatca aggagtcccg gggctggaag cactccgtga cctattcctg ctgccccga			
	accccctacc tggacatcac ctaccacttc gtcatgcagc gcctg	705		
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	<212> TYPE: PRT			
174	<213> ORGANISM: Homo sapiens			
	<400> SEQUENCE: 6			
178	Ser Glu His Glu Thr Arg Leu Val Ala Lys Leu Phe Lys Asp Tyr Ser			
179				
182	Ser Val Val Arg Pro Val Glu Asp His Arg Gln Val Val Glu Val Thr			
183				
186	Ala Gly Leu Gln Leu Ile Gln Leu Ile Asn Val Asp Glu Val Asn Gln			
187	35 40 45			
190	Ile Val Thr Thr Asn Val Arg Leu Lys Gln Gly Asp Met Val Asp Leu			
191				
194	Pro Arg Pro Ser Cys Val Thr Leu Gly Val Pro Leu Phe Ser His Leu			
195				
198	Gln Asp Glu Gln Trp Val Asp Tyr Asn Leu Lys Trp Asn Pro Asp Asp			
199				
202	Tyr Gly Gly Val Lys Lys Ile His Ile Pro Ser Glu Lys Ile Trp Arg			
203	100 105 110			
206	Pro Asp Leu Val Leu Tyr Asn Asn Ala Asp Gly Asp Phe Ala Ile Val			
207	115 120 125			
210	Lys Phe Thr Lys Val Leu Leu Gln Tyr Thr Gly His Ile Thr Trp Thr			
211	130 135 140			
214	Pro Pro Ala Ile Phe Lys Ser Tyr Cys Glu Ile Ile Val Thr His Phe			
215	145 150 155 160			
218	Pro Phe Asp Glu Gln Asn Cys Ser Met Lys Leu Gly Thr Trp Thr Tyr			
219	165 170 175			
222	Asp Gly Ser Val Val Ala Ile Asn Pro Glu Ser Asp Gln Pro Asp Leu			
223				
226	Ser Asn Phe Met Glu Ser Gly Glu Trp Val Ile Lys Glu Ser Arg Gly			
227				
230	Trp Lys His Ser Val Thr Tyr Ser Cys Cys Pro Asp Thr Pro Tyr Leu			
231	210 215 220			
234	Asp Ile Thr Tyr His Phe Val Met Gln Arg Leu			
235	225 230 235			
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241	<213> ORGANISM: Homo sapiens			
	<400> SEQUENCE: 7			
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	ccagtggaag accaccgcca ggtcgtggag gtcaccgtgg gcctgcagct gatacagct			
	atcaatgtgg atgaagtaaa tcagatcgtg acaaccaatg tgcgtctgaa acagggtga			
	50 atggtagate tgccaegece cagetgegtg actttgggag tteetttgtt tteteatetg			
252	caggatgagc aatgggtgga ttacaaccta aaatggaatc cagatgacta tggcggtgt	g 300		

Input Set : A:\sequence listing.txt
Output Set: N:\CRF4\12132005\I820339B.raw

256 258 260 262 264 266 269	aaaaaaattc acatt gcagatggtg acttt atcacgtgga cacct ccctttgatg aacag gtggccatca acccg tgggtgatca aggag accccctacc tggac <210> SEQ ID NO:	tgctat tgtca tccagc catct gaactg cagca ggaaag cgacc gtcccg gggct catcac ctacc : 8	agttc acc ttaaa agc tgaag ctc agcca gac ggaag cac	caaagtgc ctactgtg gggcacct cctgagca	tcctgcagta agatcatcgt ggacctacga acttcatgga	cactggccac cacccacttt cggctctgtc gagcggggag	360 420 480 540 600 660
	<211> DENGIA: 23	30					
	<213> ORGANISM:	Home carion	~				
		-	5				
	<400> SEQUENCE:		מות ובעד	Tria Ton	Dho Isra Nam	The Cons	
	Ser Glu His Glu	_	vai Aia	_	Pne Lys Asp	_	
277		5	Dan IIia	10	Well Well Cla	15	
	Ser Val Val Arg	PIO Val Glu	25 Asp nis	Arg Gin		, vai illi	
281	Ala Clar Lau Clar	I au Tla Cla		7 V-1	30	3 Gl-	
	Ala Gly Leu Gln	Leu lie Gin		ASII VAI		. ASH GIH	
285		Ace Val Ace	40	Cln Cl.	45	Acr Lou	
289	Ile Val Thr Thr	ASII VAI AIG	rea rys	GIN GIY	60	Asp Leu	
			ton Clar	Wal Dro		Hig Lou	
292	Pro Arg Pro Ser	70	nea Gry	75	Leu Phe Sei	80	
	Gln Asp Glu Gln	· -	Tur Acn		Trn Acn Dro		
297	GIN ASP GIU GIN	85	TYL ASII	90	TIP ASII PIC	95	
	Tyr Gly Gly Val		uic Tlo		Clu ive Tle		
301	100	nys nys ite	105	FIO SEI	110		
	Pro Asp Leu Val	Leu Tur Acn		Acn Gly			
305	115 ASP Bed Val	neu Tyr Asii	120	ASP GIY	125	. IIC vai	
	Lys Phe Thr Lys	Val Len Len		Thr Gly		Trn Thr	
309	130	135	GIN TYL	IIII GIY	140	TIP IIII	
	Pro Pro Ala Ile		Tvr Cvs	Glu Tle		His Phe	
	145	150	IJI CJU	155	110 /41 1111	160	
	Pro Phe Asp Glu		Ser Met		Gly Thr Trr		
317		165	302 1.00	170	0-7 <u>F</u>	175	
	Asp Gly Ser Val		Asn Pro		Asp Gln Pro		
321	180		185		190	_	
	Ser Asn Phe Met	Glu Ser Glv		Val Ile			
325	195		200		205	5 +-1	
	Trp Lys His Ser	Val Thr Tvr		Cvs Pro		Tvr Leu	
329	210	215		-2	220	-1	
	Asp Ile Thr Tyr						
	225	230					
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	<211> LENGTH: 20						
	338 <212> TYPE: DNA						
	39 <213> ORGANISM: Artificial						
341	41 <220> FEATURE:						
342	342 <223> OTHER INFORMATION: synthetic						
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Input Set : A:\sequence listing.txt
Output Set: N:\CRF4\12132005\I820339B.raw

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350	<212> TYPE: DNA	
351	<213> ORGANISM: Artificial	
353	<220> FEATURE:	
354	<223> OTHER INFORMATION: synthetic	
356	<400> SEQUENCE: 10	
357	cggaattcca ggcgctgcat gac	23
360	<210> SEQ ID NO: 11	
361	<211> LENGTH: 26	
362	<212> TYPE: DNA	
363	<213> ORGANISM: Artificial	
365	<220> FEATURE:	
366	<223> OTHER INFORMATION: synthetic	
368	<400> SEQUENCE: 11	
	cggaattctg gaggtgtcca cgtgat	26
372	<210> SEQ ID NO: 12	
373	<211> LENGTH: 23	
	<212> TYPE: DNA	
	<213> ORGANISM: Artificial	
	<220> FEATURE:	
	<223> OTHER INFORMATION: synthetic	
	<400> SEQUENCE: 12	
	ccggatccgc catctttaaa agc	23
	<210> SEQ ID NO: 13	
	<211> LENGTH: 25	
	<212> TYPE: DNA	
	<213> ORGANISM: Artificial	
	<220> FEATURE:	
	<223> OTHER INFORMATION: synthetic	
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	ggccatgggc tccgaacatg agacc <210> SEQ ID NO: 14	23
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	<211> LENGTH: 29 <212> TYPE: DNA	
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	<213> ORGANISM: Artificial	
	<220> FEATURE:	
	<223> OTHER INFORMATION: synthetic	
	<400> SEQUENCE: 15	
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Input Set : A:\sequence listing.txt

Output Set: N:\CRF4\12132005\1820339B.raw

## Invalid <213> Response:

Use of "Artificial" only as "<213> Organism" response is incomplete, per 1.823(b) of New Sequence Rules. Valid response is Artificial Sequence.

Seq#:9,10,11,12,13,14,15,16,17,18,19,20,21,22,23,24,25,26,27,28,29,30,31,32

VERIFICATION SUMMARYDATE: 12/13/2005PATENT APPLICATION: US/09/820,339BTIME: 14:01:38

Input Set : A:\sequence listing.txt

Output Set: N:\CRF4\12132005\1820339B.raw

L:12 M:270 C: Current Application Number differs, Replaced Current Application No

L:12 M:271 C: Current Filing Date differs, Replaced Current Filing Date